

SEQUENCE LISTING

<1<110> IRD

<120> BACTERIAL STRAINS OF GENUS EXIGOBACTERIUM, CULTURE METHOD AND USES

<130> 1721-94

<140> 10/538,715

<141> 2006-06-14

<150> PCT/FR03/003665

<151> 2003-12-10

<150> FR 02 15 865

<151> 2002-12-13

<160> 1

<170> PatentIn version 3.1

<210> 1

<211> 1510

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Exiguobacterium acetylium

<220>

<221> misc_feature

<222> (1117)..(1117)

<223> unknown

<400> 1

gcgtgcctaa tacatgcaag tcgagcgcag gaagccgtct gaacccttcg gggggacgac

120 ggtggaatga gcggcggacg ggtgagtaac acgtaaagaa cctgcccata ggtctgggat 180 aaccacgaga aatcggggct aataccggat gtgtcatcgg accgcatggt ccgctgatga aaggcgctcc ggcgtcgccc atggatggct ttgcggtgca ttagctagtt ggtggggtaa 240 cggcccacca aggcgacgat gcatagccga cctgagaggg tgatcggcca cactgggact 300 gagacacggc ccagactcct acgggaggca gcagtaggga atcttccaca atggacgaaa 360 420 gtctgatgga gcaacgccgc gtgaacgatg aaggctttcg ggtcgtaaag ttctgttgta 480 agggaagaac aagtgccgca ggcaatggcg gcaccttgac ggtaccttgc gagaaagcca 540 cggctaacta cgtgccagca gccgcggtaa tacgtaggtg gcaagcgttg tccggaatta 600 ttgggcgtaa agcgcgcgca ggcggcctct taagtctgat gtgaaagccc ccggctcaac 660 cggggagggc cattggaaac tgggaggctt gagtatagga gagaagagtg gaattccacg 720 tgtagcggtg aaatgcgtag agatgtggag gaacaccagt ggcgaaggcg actctttggc ctataactga cgctgaggct gcgaaagcgt ggggagcaaa caggattaga taccctggta 780 gtccacgccg taaacgatga gtgctaggtg ttggagggtt tccgcccttc agtgctgaag 840 ctaacgcatt aagcactccg cctggggagt acggtcgcaa ggctgaaact caaaggaatt 900 960 gacggggacc cgcacaagcg gtggagcatg tggtttaatt cgaagcaacg cgaagaacct 1020 taccaactct tgacatcccc ctgaccggta cagagatgta ccttcccctt cgggggcagg 1080 1140 caacgagcgc aaccettgtc cttagttgcc agcattnagt tgggcactct agggagactg ccggtgacaa accggaggaa ggtggggatg acgtcaaatc atcatgcccc ttatgagttg 1200 1260 ggctacacac gtgctacaat ggacggtaca aagggcagcg aagccgcgag gtggagccaa tcccagaaag ccgttctcag ttcggattgc aggctgcaac tcgcctgcat gaagtcggaa 1320 togotagtaa togoaggtoa goatactgog gtgaatacgt tocogggtot tgtacacaco 1380 gcccgtcaca ccacgagagt ttgcaacacc cgaagtcggt gaggtaaccg taaggagcca 1440 gccgccgaag gtggggcaga tgattggggt gaagtcgtaa caaggtagcc gtatcggaag 1500 1510 gtgcggctga

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Advantageously, at least 70% of the genome of the strains of the invention is capable of hybridizing with the DNA of the deposited strain.

The invention is directed in particular toward the bacterial strains defined above, characterized by the sequence SEQ ID No. 1 of the 16S rRNA:

GCGTGCCTAATACATGCAAGTCGAGCGCAGGAAGCCGTCTGAACCCTTCGGGGGGACGACGGTGGAATGA GCGGCGGACG

GGTGAGTAACACGTAAAGAACCTGCCCATAGGTCTGGGATAACCACGAGAAATCGGGGCTAATACCGGATGTCATCGG

 $\tt CGGCCCACCAAGGCGACGATGCATAGCCGACCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACGGCCCAGACTCCT$

ACGGGAGGCAGCAGTAGGGAATCTTCCACAATGGACGAAAGTCTGATGGAGCAACGCCGCGTGAACGATGAAGGCTTTCG

 ${\tt GGTCGTAAAGTTCTGTTGTAAGGGAAGAACAAGTGCCGCAGGCAATGGCGGCACCTTGACGGTACCTTGC}\\ {\tt GAGAAAGCCA}\\$

CGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGTGGCAAGCGTTGTCCGGAATTATTGGGCGTAA AGCGCGCGCA

 $\tt GGCGGCCTCTTAAGTCTGATGTGAAAGCCCCCGGCTCAACCGGGGAGGGCCATTGGAAACTGGGAGGCTTGGATATAGGA$

GAGAAGAGTGGAATTCCACGTGTAGCGGTGAAATGCGTAGAGATGTGGAGGAACACCAGTGGCGAAGGCGACTCTTTGGC

CTATAACTGACGCTGAGGCTGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCCTGGTAGTCCACGCCG
TAAACGATGA

 ${\tt GTGCTAGGTGTTGGAGGGTTTCCGCCCTTCAGTGCTGAAGCTAACGCATTAAGCACTCCGCCTGGGGAGTACGGTCGCAA}$

 ${\tt GGCTGAAACTCAAAGGAATTGACGGGGACCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGAAGCAACGCGGAAGAACCT}$

TGTCGTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTCCTTAGTTGCC AGCATTnAGT

TGGGCACTCTAGGGAGACTGCCGGTGACAAACCGGAGGAAGGTGGGGATGACGTCAAATCATCATGCCCC TTATGAGTTG

 $\tt GGCTACACGTGCTACAATGGACGGTACAAAGGGCAGCGAAGCCGCGAGGTGGAGCCAATCCCAGAAAGCCGTTCTCAG$

TTCGGATTGCAGGCTGCAACTCGCCTGCATGAAGTCGGAATCGCTAGTAATCGCAGGTCAGCATACTGCGGTGAATACGT

 ${\tt TCCCGGGTCTTGTACACACCGCCGTCACACCACGAGGTTTGCAACACCCGAAGTCGGTGAGGTAACCGTAAGGAGCCA}$

GCCGCCGAAGGTGGGCCAGATGATTGGGGTGAAGTCGTAACAAGGTAGCCGTATCGGAAGGTGCGGCTGA

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or a sequence having more than 97% similarity with SEO ID No. 1.

According to another aspect, these strains are 15 characterized in that they are thermoresistant, saccharolytic and amylolytic and/or in that they are capable of producing lactate.